

SEQUENCE LISTING

| <110> Advisys | s, Inc. |
|---------------|---------|
|---------------|---------|

<120> Codon optimized Synthetic Plasmid

<130> 108328.00146

<160> 43

<170> PatentIn version 3.3

<210> 1

<211> 3534

<212> DNA

<213> artificial sequence

<220>

<223> Plasmid vector having an analog GHRH sequence.

<400> 1

gttgtaaaac gacggccagt gaattgtaat acgactcact atagggcgaa ttggagctcc 60 120 accgcggtgg cggccgtccg cectcggcac catcctcacg acacccaaat atggcgacgg gtgaggaatg gtggggagtt atttttagag cggtgaggaa ggtgggcagg cagcaggtgt 180 240 tggcgctcta aaaataactc ccgggagtta tttttagagc ggaggaatgg tggacaccca 300 aatatggcga cggttcctca cccgtcgcca tatttgggtg tccgccctcg gccggggccg 360 catteetggg ggeeggegg tgeteeegee egeetegata aaaggeteeg gggeeggegg cggcccacga gctacccgga ggagcgggag gcgccaagct ctagaactag tggatcccaa 420 ggcccaactc cccgaaccac tcagggtcct gtggacagct cacctagctg ccatggtgct 480 ctgggtgttc ttctttgtga tcctcaccct cagcaacagc tcccactgct ccccacctcc 540

600 ccctttgacc ctcaggatgc ggcggcacgt agatgccatc ttcaccaaca gctaccggaa ggtgctggcc cagctgtccg cccgcaagct gctccaggac atcctgaaca ggcagcaggg 660 720 agagaggaac caagagcaag gagcataatg actgcaggaa ttcgatatca agcttatcgg ggtggcatcc ctgtgacccc tccccagtgc ctctcctggc cctggaagtt gccactccag 780 840 tgcccaccag ccttgtccta ataaaattaa gttgcatcat tttgtctgac taggtgtcct tctataatat tatggggtgg aggggggtgg tatggagcaa ggggcaagtt gggaagacaa 900 960 cctgtagggc ctgcggggtc tattgggaac caagctggag tgcagtggca caatcttggc 1020 tcactgcaat ctccgcctcc tgggttcaag cgattctcct gcctcagcct cccgagttgt 1080 tgggattcca ggcatgcatg accaggctca gctaattttt gtttttttgg tagagacggg gtttcaccat attggccagg ctggtctcca actcctaatc tcaggtgatc tacccacctt 1140 1200 1260 ttttaaaata actataccag caggaggacg tccagacaca gcataggcta cctggccatg 1320 cccaaccggt gggacatttg agttgcttgc ttggcactgt cctctcatgc gttgggtcca 1380 ctcagtagat gcctgttgaa ttcgataccg tcgacctcga gggggggccc ggtaccagct 1440 tttgttccct ttagtgaggg ttaatttcga gcttggcgta atcatggtca tagctgtttc 1500 ctgtgtgaaa ttgttatccg ctcacaattc cacacaacat acgagccgga agcataaagt 1560 gtaaagcctg gggtgcctaa tgagtgagct aactcacatt aattgcgttg cgctcactgc 1620 ccgctttcca gtcgggaaac ctgtcgtgcc agctgcatta atgaatcggc caacgcgcgg ggagaggegg tttgegtatt gggegetett eegetteete geteaetgae tegetgeget 1680

1740 cggtcgttcg gctgcggcga gcggtatcag ctcactcaaa ggcggtaata cggttatcca 1800 cagaatcagg ggataacgca ggaaagaaca tgtgagcaaa aggccagcaa aaggccagga 1860 accgtaaaaa ggccgcgttg ctggcgtttt tccataggct ccgccccct gacgagcatc 1920 acaaaaatcg acgctcaagt cagaggtggc gaaacccgac aggactataa agataccagg cgtttccccc tggaagctcc ctcgtgcgct ctcctgttcc gaccctgccg cttaccggat 1980 2040 acctgtccgc ctttctccct tcgggaagcg tggcgctttc tcatagctca cgctgtaggt 2100 atctcagttc ggtgtaggtc gttcgctcca agctgggctg tgtgcacgaa ccccccgttc 2160 agcccgaccg ctgcgcctta tccggtaact atcgtcttga gtccaacccg gtaagacacg 2220 acttategee actggeagea gecactggta acaggattag cagagegagg tatgtaggeg 2280 gtgctacaga gttcttgaag tggtggccta actacggcta cactagaaga acagtatttg gtatctgcgc tctgctgaag ccagttacct tcggaaaaag agttggtagc tcttgatccg 2340 gcaaacaaac caccgctggt agcggtggtt tttttgtttg caagcagcag attacgcgca 2400 gaaaaaaagg atctcaagaa gatcctttga tcttttctac ggggtctgac gctcagaaga 2460 2520 actegteaag aaggegatag aaggegatge getgegaate gggageggeg atacegtaaa gcacgaggaa gcggtcagcc cattcgccgc caagctcttc agcaatatca cgggtagcca 2580 2640 acgctatgtc ctgatagcgg tccgccacac ccagccggcc acagtcgatg aatccagaaa 2700 ageggeeatt ttecaceatg atatteggea ageaggeate geeatgggte aegaegagat 2760 cetegeegte gggeatgege geettgagee tggegaacag tteggetgge gegageeeet 2820 gatgetette gtecagatea teetgatega caagacegge ttecateega gtaegtgete

2880 gctcgatgcg atgtttcgct tggtggtcga atgggcaggt agccggatca agcgtatgca 2940 gccgccgcat tgcatcagcc atgatggata ctttctcggc aggagcaagg tgagatgaca 3000 qqaqatcctg ccccggcact tcgcccaata gcagccagtc ccttcccgct tcagtgacaa 3060 cgtcgagcac agctgcgcaa ggaacgcccg tcgtggccag ccacgatagc cgcgctgcct cgtcctgcag ttcattcagg gcaccggaca ggtcggtctt gacaaaaaga accgggcgcc 3120 3180 cctgcgctga cagccggaac acggcggcat cagagcagcc gattgtctgt tgtgcccagt catageegaa tageetetee acceaagegg eeggagaace tgegtgeaat eeatettgtt 3240 3300 caatcatgcg aaacgatcct catcctgtct cttgatcaga tcttgatccc ctgcgccatc 3360 agateettgg eggeaagaaa geeateeagt ttaetttgea gggetteeea acettaeeag 3420 agggegeece agetggeaat teeggttege ttgetgteea taaaacegee cagtetagea actgttggga agggcgatcg gtgcgggcct cttcgctatt acgccagctg gcgaaagggg 3480 gatgtgctgc aaggcgatta agttgggtaa cgccagggtt ttcccagtca cgac 3534

<210> 2

<211> 2739

<212> DNA

<213> artificial sequence

<220>

<223> Optimized vector having an analog GHRH sequence.

<400> 2

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180 gttggcgctc taaaaataac tcccgggagt tatttttaga gcggaggaat ggtggacacc 240 caaatatggc gacggttcct cacccgtcgc catatttggg tgtccgccct cggccggggc 300 cgcattcctg ggggccgggc ggtgctcccg cccgcctcga taaaaggctc cggggccggc 360 ggcggcccac gagctacccg gaggagcggg aggcgccaag cggatcccaa ggcccaactc 420 cccgaaccac tcagggtcct gtggacagct cacctagctg ccatggtgct ctgggtgttc 480 ttctttgtga tcctcaccct cagcaacagc tcccactgct ccccacctcc ccctttgacc ctcaggatgc ggcggtatgc agatgccatc ttcaccaaca gctaccggaa ggtgctgggc 540 600 cagctgtccg cccgcaagct gctccaggac atcatgagca ggcagcaggg agagaggaac 660 caagagcaag gagcataatg actgcaggaa ttcgatatca agcttatcgg ggtggcatcc 720 ctgtgacccc tccccagtgc ctctcctggc cctggaagtt gccactccag tgcccaccag ccttgtccta ataaaattaa gttgcatcat tttgtctgac taggtgtcct tctataatat 780 tatggggtgg agggggtgg tatggagcaa ggggcaagtt gggaagacaa cctgtagggc 840 900 tcgaggggg gcccggtacc agcttttgtt ccctttagtg agggttaatt tcgagcttgg 960 tetteegett cetegeteae tgaetegetg egeteggteg tteggetgeg gegageggta 1020 tcagctcact caaaggcggt aatacggtta tccacagaat caggggataa cgcaggaaag 1080 aacatgtgag caaaaggcca gcaaaaggcc aggaaccgta aaaaggccgc gttgctggcg tttttccata ggctccgccc ccctgacgag catcacaaaa atcgacgctc aagtcagagg 1140 tggcgaaacc cgacaggact ataaagatac caggcgtttc cccctggaag ctccctcgtg 1200 cgctctcctg ttccgaccct gccgcttacc ggatacctgt ccgcctttct cccttcggga 1260

1320 agcqtqqcgc tttctcatag ctcacgctgt aggtatctca gttcggtgta ggtcgttcgc tccaagetgg getgtgtgca egaaceeece gttcageeeg acegetgege ettateeggt 1380 1440 aactatcgtc ttgagtccaa cccggtaaga cacgacttat cgccactggc agcagccact 1500 ggtaacagga ttagcagagc gaggtatgta ggcggtgcta cagagttctt gaagtggtgg cctaactacg gctacactag aagaacagta tttggtatct gcgctctgct gaagccagtt 1560 1620 accttcggaa aaagagttgg tagctcttga tccggcaaac aaaccaccgc tggtagcggt ggtttttttg tttgcaagca gcagattacg cgcagaaaaa aaggatctca agaagatcct 1680 1740 ttgatctttt ctacggggtc tgacgctcag ctagcgctca gaagaactcg tcaagaaggc gatagaaggc gatgcgctgc gaatcgggag cggcgatacc gtaaagcacg aggaagcggt 1800 cageceatte geogecaage tetteageaa tateaegggt agecaaeget atgteetgat 1860 ageggteege cacacceage eggeeacagt egatgaatee agaaaagegg ecatttteea 1920 1980 ccatgatatt cggcaagcag gcatcgccat gagtcacgac gagatcctcg ccgtcgggca 2040 tgcgcgcctt gagcctggcg aacagttcgg ctggcgcgag cccctgatgc tcttcgtcca gatcatcctg atcgacaaga ccggcttcca tccgagtacg tgctcgctcg atgcgatgtt 2100 2160 tegettggtg gtegaatggg caggtageeg gateaagegt atgeageege egeattgeat 2220 cagccatgat ggatactttc tcggcaggag caaggtgaga tgacaggaga tcctgccccg 2280 gcacttegee caatageage cagtecette eegetteagt gacaaegteg ageacagetg 2340 cgcaaggaac gcccgtcgtg gccagccacg atagccgcgc tgcctcgtcc tgcagttcat 2400 tcagggcacc ggacaggtcg gtcttgacaa aaagaaccgg gcgcccctgc gctgacagcc

ggaacacggc ggcatcagag cagccgattg tctgttgtgc ccagtcatag ccgaatagcc 2460

tctccaccca agcggccgga gaacctgcgt gcaatccatc ttgttcaatc atgcgaaacg 2520

atcctcatcc tgtctcttga tcagatcttg atcccctgcg ccatcagatc cttggcggca 2580

agaaagccat ccagtttact ttgcagggct tcccaacctt accagaggc gccccagctg 2640

gcaattccgg ttcgcttgct gtccataaaa ccgcccagtc tagcaactgt tgggaagggc 2700

gatcgtgtaa tacgactcac tatagggcga attggagct 2739

<210> 3

<211> 795

<212> DNA

<213> artificial sequence

<220>

<223> Nucleic acid sequence for the antibiotic resistance gene kanamycin.

<400> 3

atgattgaac aagatggatt gcacgcaggt tctccggccg cttgggtgga gaggctattc 60

ggctatgact gggcacaaca gacaatcggc tgctctgatg ccgccgtgtt ccggctgtca 120

gcgcaggggc gcccggttct ttttgtcaag accgacctgt ccggtgccct gaatgaactg 180

caggacgagg cagcgcggct atcgtggctg gccacgacgg gcgttccttg cgcagctgtg 240

ctcgacgttg tcactgaagc gggaagggac tggctgctat tgggcgaagt gccggggcag 300

gatctcctgt catctcacct tgctcctgcc gagaaagtat ccatcatggc tgatgcaatg 360

cggcggctgc atacgcttga tccggctacc tgcccattcg accaccaagc gaaacatcgc 420

| atcgagcgag | cacgtactcg | gatggaagcc | ggtcttgtcg | atcaggatga | tctggacgaa | 480 |
|------------|------------|------------|------------|------------|------------|-----|
| gagcatcagg | ggctcgcgcc | agccgaactg | ttcgccaggc | tcaaggcgcg | catgcccgac | 540 |
| ggcgaggatc | tcgtcgtgac | tcatggcgat | gcctgcttgc | cgaatatcat | ggtggaaaat | 600 |
| ggccgctttt | ctggattcat | cgactgtggc | cggctgggtg | tggcggaccg | ctatcaggac | 660 |
| atagcgttgg | ctacccgtga | tattgctgaa | gagettggeg | gcgaatgggc | tgaccgcttc | 720 |
| ctcgtgcttt | acggtatcgc | cgctcccgat | tcgcagcgca | tcgccttcta | tcgccttctt | 780 |
| gacgagttct | tctga | | | | | 795 |

<210> 4

<211> 219

<212> DNA

<213> artificial sequence

<220>

<223> Sequence for an analog porcine GHRH sequence.

<400> 4

atggtgctct gggtgttctt ctttgtgatc ctcaccctca gcaacagctc ccactgctcc 60

ccacctcccc ctttgaccct caggatgcgg cggcacgtag atgccatctt caccaacagc 120

taccggaagg tgctggccca gctgtccgcc cgcaagctgc tccaggacat cctgaacagg 180

cagcagggag agaggaacca agagcaagga gcataatga 219

<210> 5

<211> 246

<212> DNA

<213> artificial sequence

| _ | 2 | 2 | n | |
|---|---|---|---|--|

<223> Sequence for an analog mouse GHRH sequence.

<400> 5

gccatggtgc tctgggtgct ctttgtgatc ctcatcctca ccagcggcag ccactgcagc 60
ctgcctccca gccctccctt caggatgcag aggcacgtgg acgccatctt caccaccaac 120
tacaggaagc tgctgagcca gctgtacgcc aggaaggtga tccaggacat catgaacaag 180
cagggcgaga ggatccagga gcagagggcc aggctgagct gataagcttg cgatgagttc 240
ttctaa 246

<210> 6

<211> 234

<212> DNA

<213> artificial sequence

<220>

<223> Sequence for an analog rat GHRH sequence.

<400> 6

gecatggee tgtgggtgtt ettegtgetg etgaceetga eeageggaag eeactgeage 60
etgeeteeca geeeteeett eagggtgege eggeaegeeg aegeeatett eaceageage 120
tacaggagga teetgggeea getgtaeget aggaagetee tgeaegagat eatgaaeagg 180
eageagggeg agaggaaeca ggageagagg ageaggttea aetgataage ttge 234

<210> 7

<211> 225

<212> DNA

<213> artificial sequence

| | _ | _ | _ | |
|---|---|---|---|--|
| _ | า | 2 | Λ | |
| | | | | |

<223> Sequence for an analog bovine GHRH sequence.

<400> 7

gccatggtgc tgtgggtgtt cttcctggtg accctgaccc tgagcagcgg ctcccacggc 60

tecctgccct cccagcctct gcgcatccct cgctacgccg acgccatctt caccaacagc 120

taccgcaagg tgctcggcca gctcagcgcc cgcaagctcc tgcaggacat catgaaccgg 180

cagcagggcg agcgcaacca ggagcaggga gcctgataag cttgc 225

<210> 8

<211> 225

<212> DNA

<213> artificial sequence

<220>

<223> Sequence for an analog ovine GHRH sequence.

<400> 8

gccatggtgc tgtgggtgtt cttcctggtg accctgaccc tgagcagcgg aagccacggc 60
agcctgccca gccagcccct gaggatccct aggtacgccg acgccatctt caccaacagc 120
tacaggaaga tcctgggcca gctgagcgct aggaagctcc tgcaggacat catgaacagg 180
cagcagggcg agaggaacca ggagcagggc gcctgataag cttgc 225

<210> 9

<211> 246

<212> DNA

<213> artificial sequence

<220>

<223> Sequence for an analog chicken GHRH sequence.

| <400> | 9 | | | | | | |
|---------|-----------|--------------|------------|-------------|--------------|--------------|-----|
| gccatg | gtgc | tctgggtgct | ctttgtgatc | ctcatcctca | ccagcggcag | ccactgcagc | 60 |
| ctgcct | ccca | gccctccctt | caggatgcag | aggcacgtgg | acgccatctt | caccaccaac | 120 |
| tacagga | aagc | tgctgagcca | gctgtacgcc | aggaaggtga | tccaggacat | catgaacaag | 180 |
| cagggc | gaga | ggatccagga | gcagagggcc | aggctgagct | gataagcttg | cgatgagttc | 240 |
| ttctaa | | | | | | | 246 |
| | | | | | | | |
| <210> | 10 | | | | | | |
| <211> | 190 | | | | | | |
| <212> | DNA | | | | | | |
| <213> | art: | ificial seq | uence | | | | |
| <220> | | | | | | | |
| <223> | Nuc: | leic acid s | equence of | human growt | th hormone p | ooly A tail. | |
| <400> | 10 | | | | | | |
| gggtgg | catc | cctgtgaccc | ctccccagtg | cctctcctgg | ccctggaagt | tgccactcca | 60 |
| gtgccca | acca | gccttgtcct | aataaaatta | agttgcatca | ttttgtctga | ctaggtgtcc | 120 |
| ttctata | aata | ttatggggtg | gaggggggtg | gtatggagca | aggggcaagt | tgggaagaca | 180 |
| acctgta | aggg | | | | | | 190 |
| -210- | 11 | | | | | | |
| <210> | 11 | | | | | | |
| <211> | 55 DVA | | | | | | |
| <212> | DNA | . 6 | | | | | |
| <213> | arti | ificial sequ | uence | | | | |
| <220> | | | | | | | |

<223> Nucleic acid sequence of human growth hormone 5' untranslated

region

| <400> | 11 | | | | | | |
|---------|------|--------------|--------------|-------------|--------------|--------------|------|
| caaggcc | caa | ctccccgaac | cactcagggt | cctgtggaca | gctcacctag | ctgcc | 55 |
| | | | | | | | |
| | | | | | | | |
| <210> | 12 | | | | | | |
| <211> | 782 | | | | | | |
| <212> | DNA | | | | | | |
| <213> | art: | ificial sequ | uence | | | | |
| | | | | | | | |
| <220> | | | | | | | |
| <223> | Nuc. | leic acid se | equence of a | a plasmid p | JC-18 origin | n of replica | iton |
| | | | | | | | |
| <400> | 12 | | | | | | |
| tcttccg | ctt | cctcgctcac | tgactcgctg | cgctcggtcg | ttcggctgcg | gcgagcggta | 60 |
| | | | | | | | |
| tcagctc | act | caaaggcggt | aatacggtta | tccacagaat | caggggataa | cgcaggaaag | 120 |
| | | | | | | | |
| aacatgt | gag | caaaaggcca | gcaaaaggcc | aggaaccgta | aaaaggccgc | gttgctggcg | 180 |
| | | | | | | | |
| tttttcc | ata | ggctccgccc | ccctgacgag | catcacaaaa | atcgacgctc | aagtcagagg | 240 |
| | | | | | | | |
| tggcgaa | acc | cgacaggact | ataaagatac | caggcgtttc | cccctggaag | ctccctcgtg | 300 |
| | | | • | | | | |
| cgctctc | ctg | ttccgaccct | gccgcttacc | ggatacctgt | ccgcctttct | cccttcggga | 360 |
| | | | | | | | |
| agcgtgg | cgc | tttctcatag | ctcacgctgt | aggtatctca | gttcggtgta | ggtcgttcgc | 420 |
| | | | | | | | |
| tccaago | tgg | gctgtgtgca | cgaacccccc | gttcagcccg | accgctgcgc | cttatccggt | 480 |
| | | | | | | | |
| aactatc | gtc | ttgagtccaa | cccggtaaga | cacgacttat | cgccactggc | agcagccact | 540 |
| | | | | | | | |
| ggtaaca | gga | ttagcagagc | gaggtatgta | ggcggtgcta | cagagttctt | gaagtggtgg | 600 |
| | | | | | | | |
| cctaact | acg | gctacactag | aaggacagta | tttggtatct | gcgctctgct | gaagccagtt | 660 |
| | | | | | | | |
| accttcg | gaa | aaagagttgg | tagctcttga | tccggcaaac | aaaccaccgc | tggtagcggt | 720 |

| ggtttt | tttg tttgcaagca gcagattacg cgcagaaaaa aaggatctca agaagatcct | 780 |
|--------|---|-----|
| tt | | 782 |
| | | |
| | | |
| <210> | 13 | |
| <211> | 5 | |
| <212> | DNA | |
| <213> | artificial sequence | |
| | | |
| <220> | | |
| <223> | This is a NEO ribosomal binding site | |
| | | |
| <400> | 13 | - |
| tcctc | | 5 |
| | | |
| <210> | 14 | |
| <211> | 29 | |
| <212> | DNA | |
| <213> | artificial sequence | |
| | | |
| <220> | | |
| <223> | Nucleic acid sequence of a prokaryotic PNEO promoter. | |
| | | |
| <400> | 14 | |
| acctta | ccag agggcgcccc agctggcaa | 29 |
| | | |
| | | |
| <210> | 15 | |
| <211> | 323 | |
| <212> | DNA | |
| <213> | artificial sequence | |
| <220> | | |
| <223> | Nucleic acid sequence of a eukaryotic promoter c5-12. | |
| 12237 | native and bequence of a canalyoute promoter to 12. | |

| 15 | | | | | | |
|------|------------|--|---|--|---|--|
| ccg | ccctcggcac | catcctcacg | acacccaaat | atggcgacgg | gtgaggaatg | 60 |
| gtt | atttttagag | cggtgaggaa | ggtgggcagg | cagcaggtgt | tggcgctcta | 120 |
| ctc | ccgggagtta | tttttagagc | ggaggaatgg | tggacaccca | aatatggcga | 180 |
| tca | cccgtcgcca | tatttgggtg | teegeeeteg | gccggggccg | cattcctggg | 240 |
| ıcgg | tgctcccgcc | cgcctcgata | aaaggctccg | gggccggcgg | cggcccacga | 300 |
| gga | ggagcgggag | gcg | | | | 323 |
| | eccg | eccg ccctcggcac gtt atttttagag ctc ccgggagtta ctca cccgtcgcca | eccg ccctcggcac catcctcacg gtt atttttagag cggtgaggaa ectc ccgggagtta tttttagagc etca cccgtcgcca tatttgggtg | ccg ccctcggcac catcctcacg acacccaaat gtt atttttagag cggtgaggaa ggtgggcagg ctc ccgggagtta tttttagagc ggaggaatgg ctca cccgtcgcca tatttgggtg tccgccctcg gcgg tgctcccgcc cgcctcgata aaaggctccg | ccg ccctcggcac catcctcacg acacccaaat atggcgacgg ggt attttagag cggtgaggaa ggtgggcagg cagcaggtgt ctc ccgggagtta tttttagagc ggaggaatgg tggacaccca etca cccgtcgcca tatttgggtg tccgccctcg gccggggccg ggg tgctcccgcc cgcctcgata aaaggctccg gggccggcgg | ccg ccctcggcac catcctcacg acacccaaat atggcgacgg gtgaggaatg ggt atttttagag cggtgaggaa ggtgggcagg cagcaggtgt tggcgctcta ccc ccgggagtta tttttagagc ggaggaatgg tggacaccca aatatggcga cccggtcgcca tatttgggtg tccgcctcg gccggggccg cattcctggg cgg tgctcccgcc cgcctcgata aaaggctccg gggccggcgg cggcccacga |

<210> 16

<211> 210

<212> DNA

<213> artificial sequence

<220>

<223> Optimized nucleic acid sequence of a human growth hormone poly A tail

<400> 16

ttatcggggt ggcatccctg tgacccctcc ccagtgcctc tcctggccct ggaagttgcc 60
actccagtgc ccaccagcct tgtcctaata aaattaagtt gcatcatttt gtctgactag 120
gtgtccttct ataatattat ggggtggagg ggggtggtat ggagcaaggg gcaagttggg 180
aagacaacct gtagggctcg aggggggcc 210

<210> 17

<211> 2722

<212> DNA

<213> artificial sequence

Plasmid vector having a codon optimized mouse GHRH sequence <223> <400> 17 60 ccaccgcggt ggcggccgtc cgccctcggc accatcctca cgacacccaa atatggcgac 120 gggtgaggaa tggtggggag ttatttttag agcggtgagg aaggtgggca ggcagcaggt 180 gttggcgctc taadaataac tcccgggagt tatttttaga gcggaggaat ggtggacacc caaatatggc gacggttcct cacccgtcgc catatttggg tgtccgccct cggccggggc 240 300 cgcattcctg ggggccgggc ggtgctcccg cccgctcga taaaaggctc cggggccggc 360 ggeggeceae gagetaeeeg gaggageggg aggegecaag eggateeeaa ggeecaaete cccgaaccac tcagggtcct gtggacagct cacctagctg ccatggtgct ctgggtgctc 420 480 tttgtgatcc tcatcctcac cageggcagc cactgcagcc tgcctcccag ccctcccttc aggatgcaga ggcacgtgga cgccatcttc accaccaact acaggaagct gctgagccag 540 600 ctgtacgcca ggaaggtgat ccaggacatc atgaacaagc agggcgagag gatccaggag cagagggcca ggctgagctg ataagcttat cggggtggca tccctgtgac ccctcccag 660 720 tgcctctcct ggccctggaa gttgccactc cagtgcccac cagccttgtc ctaataaaat taagttgcat cattttgtct gactaggtgt ccttctataa tattatgggg tggaggggg 780 840 tggtatggag caaggggcaa gttgggaaga caacctgtag ggctcgaggg ggggcccggt accagetttt gtteeettta gtgagggtta atttegaget tggtetteeg etteeteget 900 960 cactgacteg etgegetegg tegttegget geggegageg gtateagete acteaaagge 1020 ggtaatacgg ttatccacag aatcagggga taacgcagga aagaacatgt gagcaaaagg

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2220 ttctcggcag gagcaaggtg agatgacagg agatcctgcc ccggcacttc gcccaatagc agccagtccc ttcccgcttc agtgacaacg tcgagcacag ctgcgcaagg aacgcccgtc 2280 2340 gtggccagcc acgatagccg cgctgcctcg tcctgcagtt cattcagggc accggacagg 2400 tcggtcttga caaaaagaac cgggcgcccc tgcgctgaca gccggaacac ggcggcatca gagcagccga ttgtctgttg tgcccagtca tagccgaata gcctctccac ccaagcggcc 2460 ggagaacctg cgtgcaatcc atcttgttca atcatgcgaa acgatcctca tcctgtctct 2520 tgatcagatc ttgatcccct gcgccatcag atccttggcg gcaagaaagc catccagttt 2580 2640 actttgcagg gcttcccaac cttaccagag ggcgccccag ctggcaattc cggttcgctt 2700 . gctgtccata aaaccgccca gtctagcaac tgttgggaag ggcgatcgtg taatacgact 2722 cactataggg cgaattggag ct

<210> 18

<211> 2725

<212> DNA

<213> artificial sequence

<220>

<223> Plasmid vector having a codon optimized rat GHRH sequence

<400> 18

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gttggcgctc taaaaataac tcccgggagt tatttttaga gcggaggaat ggtggacacc 180
caaatatggc gacggttcct cacccgtcgc catatttggg tgtccgcct cggccggggc 240

300 cgcattcctg ggggccgggc ggtgctcccg cccgcctcga taaaaggctc cggggccggc 360 ggcggcccac gagctacccg gaggagcggg aggcgccaag cggatcccaa ggcccaactc 420 cccgaaccac tcagggtcct gtggacaget cacctagetg ccatggccct gtgggtgttc 480 ttcgtgctgc tgaccctgac cagcggaagc cactgcagcc tgcctcccag ccctcccttc agggtgcgcc ggcacgccga cgccatcttc accagcagct acaggaggat cctgggccag 540 ctgtacgcta ggaagctcct gcacgagatc atgaacaggc agcagggcga gaggaaccag 600 gagcagagga gcaggttcaa ctgataagct tatcggggtg gcatccctgt gacccctccc 660 cagtgeetet eetggeeetg gaagttgeea etceagtgee caecageett gteetaataa 720 780 aattaagttg catcattttg tctgactagg tgtccttcta taatattatg gggtggaggg 840 gggtggtatg gagcaagggg caagttggga agacaacctg tagggctcga gggggggccc 900 ggtaccagct tttgttccct ttagtgaggg ttaatttcga gcttggtctt ccgcttcctc 960 gctcactgac tcgctgcgct cggtcgttcg gctgcggcga gcggtatcag ctcactcaaa ggcggtaata cggttatcca cagaatcagg ggataacgca ggaaagaaca tgtgagcaaa 1020 1080 aggccagcaa aaggccagga accgtaaaaa ggccgcgttg ctggcgtttt tccataggct ccgccccct gacgagcatc acaaaaatcg acgctcaagt cagaggtggc gaaacccgac 1140 1200 aggactataa agataccagg cgtttccccc tggaagctcc ctcgtgcgct ctcctgttcc gaccetgeeg ettaceggat acetgteege ettteteeet tegggaageg tggegettte 1260 1320 tcatagetca egetgtaggt ateteagtte ggtgtaggte gttegeteea agetgggetg 1380 tgtgcacgaa cccccgttc agcccgaccg ctgcgcctta tccggtaact atcgtcttga

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tettgateag atettgatee cetgegeeat cagateettg geggeaagaa agceateeag 2580

tttaetttge agggetteee aacettacea gagggegeee cagetggeaa tteeggtteg 2640

ettgetgtee ataaaacege ceagtetage aactgttggg aagggegate gtgtaatacg 2700

acteactata gggegaattg gaget 2725

<210> 19

<211> 2716

<212> DNA

<213> artificial sequence

<220>

<223> Plasmid vector having a codon optimized bovine GHRH sequence

<400> 19

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<210> 20

<211> 2716

<212> DNA

<220>

Plasmid vector having a codon optimized ovine GHRH sequence <223> <400> 20 ccaccgcggt ggcggccgtc cgccctcggc accatcctca cgacacccaa atatggcgac 60 120 gggtgaggaa tggtggggag ttatttttag agcggtgagg aaggtgggca ggcagcaggt gttggcgctc taaaaataac tcccgggagt tatttttaga gcggaggaat ggtggacacc 180 caaatatggc gacggttcct cacccgtcgc catatttggg tgtccgccct cggccggggc 240 300 cgcattcctg ggggccgggc ggtgctcccg cccgcctcga taaaaggctc cggggccggc 360 ggcggcccac gagctacccg gaggagcggg aggcgccaag cggatcccaa ggcccaactc 420 cccgaaccac tcagggtcct gtggacagct cacctagctg ccatggtgct gtgggtgttc ttcctggtga ccctgaccct gagcagcgga agccacggca gcctgcccag ccagcccctg 480 aggateceta ggtacgeega egecatette accaacaget acaggaagat eetgggeeag 540 ctgagcgcta ggaagctcct gcaggacatc atgaacaggc agcagggcga gaggaaccag 600 660 gagcagggcg cctgataagc ttatcggggt ggcatccctg tgacccctcc ccagtgcctc 720 tectggeect ggaagttgee acteeagtge ceaceageet tgteetaata aaattaagtt 780 gcatcatttt gtctgactag gtgtccttct ataatattat ggggtggagg ggggtggtat 840 ggagcaaggg gcaagttggg aagacaacct gtagggctcg agggggggcc cggtaccagc ttttgttccc tttagtgagg gttaatttcg agcttggtct tccgcttcct cgctcactga 900 ctcgctgcgc tcggtcgttc ggctgcggcg agcggtatca gctcactcaa aggcggtaat 960

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<210> 21

<211> 2725

<212> DNA

<213> artificial sequence

<220>

<223> Plasmid vector having a codon optimized chicken GHRH sequence

<400> 21

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geeggagaac etgegtgeaa tecatettgt teaateatge gaaacgatee teateetgte 2520

tettgateag atettgatee eetgegeeat eagateettg geggeaagaa ageeateeag 2580

tttaetttge agggetteee aacettacea gagggegeee eagetggeaa tteeggtteg 2640

ettgetgtee ataaaacege eeagtetage aactgttggg aagggegate gtgtaatacg 2700

acteactata gggegaattg gaget 2725

<210> 22

<211> 264

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for a coding sequence having an antibiotic resistance gene kanamycin

<400> 22

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Glu Arg Leu Phe Gly Tyr Asp Trp Ala Gln Gln Thr Ile Gly Cys Ser
20 25 30

Asp Ala Ala Val Phe Arg Leu Ser Ala Gln Gly Arg Pro Val Leu Phe 35 40 45

Val Lys Thr Asp Leu Ser Gly Ala Leu Asn Glu Leu Gln Asp Glu Ala 50 55 60

Ala Arg Leu Ser Trp Leu Ala Thr Thr Gly Val Pro Cys Ala Ala Val 65 70 75 80

Leu Asp Val Val Thr Glu Ala Gly Arg Asp Trp Leu Leu Gly Glu
85 90 95

Val Pro Gly Gln Asp Leu Leu Ser Ser His Leu Ala Pro Ala Glu Lys
100 105 110

Val Ser Ile Met Ala Asp Ala Met Arg Arg Leu His Thr Leu Asp Pro 115 120 125

Ala Thr Cys Pro Phe Asp His Gln Ala Lys His Arg Ile Glu Arg Ala 130 135 140

Arg Thr Arg Met Glu Ala Gly Leu Val Asp Gln Asp Asp Leu Asp Glu
145 150 155 160

Glu His Gln Gly Leu Ala Pro Ala Glu Leu Phe Ala Arg Leu Lys Ala 165 170 175

Arg Met Pro Asp Gly Glu Asp Leu Val Val Thr His Gly Asp Ala Cys 180 185 190

Leu Pro Asn Ile Met Val Glu Asn Gly Arg Phe Ser Gly Phe Ile Asp 195 200 205

Cys Gly Arg Leu Gly Val Ala Asp Arg Tyr Gln Asp Ile Ala Leu Ala 210 215 220 Thr Arg Asp Ile Ala Glu Glu Leu Gly Gly Glu Trp Ala Asp Arg Phe 225 230 235 240

Leu Val Leu Tyr Gly Ile Ala Ala Pro Asp Ser Gln Arg Ile Ala Phe
245 250 255

Tyr Arg Leu Leu Asp Glu Phe Phe 260

<210> 23

<211> 75

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for an analog mouse GHRH sequence

<400> 23

Ala Met Val Leu Trp Val Leu Phe Val Ile Leu Ile Leu Thr Ser Gly

1 5 10 15

Ser His Cys Ser Leu Pro Pro Ser Pro Pro Phe Arg Met Gln Arg His
20 25 30

Val Asp Ala Ile Phe Thr Thr Asn Tyr Arg Lys Leu Leu Ser Gln Leu 35 40 45

Tyr Ala Arg Lys Val Ile Gln Asp Ile Met Asn Lys Gln Gly Glu Arg 50 55 60

<210> 24 <211> 231 <212> DNA <213> artificial sequence <220> <223> Nucleic acid sequence for an original mouse GHRH sequence <400> 24 gccatggtgc tctgggtgct ctttgtgatc ctcatcctca ccagcggcag ccactgcagc 60 120 ctgcctccca gccctccctt caggatgcag aggcacgtgg acgccatctt caccaccaac tacaggaagc tgctgagcca gctgtacgcc aggaaggtga tccaggacat catgaacaag 180 cagggcgaga ggatccagga gcagagggcc aggctgagct gataagcttg c 231 <210> 25 <211> 73 <212> PRT <213> artificial sequence <220> <223> Amino acid sequence for an original mouse GHRH sequence <400> 25

75

Ile Gln Glu Gln Arg Ala Arg Leu Ser Ala Cys 70

65

10

15

Met Val Leu Trp Val Leu Phe Val Ile Leu Ile Leu Thr Ser Gly Ser

1

5

His Cys Ser Leu Pro Pro Ser Pro Pro Phe Arg Met Gln Arg His Val 20 25 30

Asp Ala Ile Phe Thr Thr Asn Tyr Arg Lys Leu Leu Ser Gln Leu Tyr 35 40 45

Ala Arg Lys Val Ile Gln Asp Ile Met Asn Lys Gln Gly Glu Arg Ile 50 55 60

Gln Glu Gln Arg Ala Arg Leu Ser Ala 65 70

<210> 26

<211> 76

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for an analog rat GHRH sequence

<400> 26

Ala Met Ala Leu Trp Val Phe Phe Val Leu Leu Thr Leu Thr Ser Gly

1 5 10 15

Ser His Cys Ser Leu Pro Pro Ser Pro Pro Phe Arg Val Arg Arg His 20 25 30

Ala Asp Ala Ile Phe Thr Ser Ser Tyr Arg Arg Ile Leu Gly Gln Leu 35 40 45

Arg Asn Gln Glu Gln Arg Ser Arg Phe Asn Ala Cys 70 <210> 27 <211> 234 <212> DNA <213> artificial sequence <220> <223> Nucleic acid sequence for an original rat GHRH sequence <400> 27 gccatggcac tctgggtgtt ctttgtgctc ctcaccctca ccagtggctc ccactgctca 60 ctgccccct cacctcctt cagggtgcgg cggcacgccg acgccatctt caccagcagc 120 tacaggagaa tcctgggcca gctgtacgcc aggaaactgc tgcacgagat catgaacagg 180 cagcagggcg agaggaacca ggagcagagg tccaggttca actgataagc ttgc 234 <210> 28 <211> 74 <212> PRT <213> artificial sequence <220> <223> Amino acid sequence for an original rat GHRH sequence <400> 28 Met Ala Leu Trp Val Phe Phe Val Leu Leu Thr Leu Thr Ser Gly Ser

Tyr Ala Arg Lys Leu Leu His Glu Ile Met Asn Arg Gln Gln Gly Glu

60

55

50

15

10

5

1

His Cys Ser Leu Pro Pro Ser Pro Pro Phe Arg Val Arg Arg His Ala 20 25 30

Asp Ala Ile Phe Thr Ser Ser Tyr Arg Arg Ile Leu Gly Gln Leu Tyr 35 40 45

Ala Arg Lys Leu Leu His Glu Ile Met Asn Arg Gln Gln Gly Glu Arg 50 55 60

Asn Gln Glu Gln Arg Ser Arg Phe Asn Ala 65 70

<210> 29

<211> 73

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for an analog bovine GHRH sequence

<400> 29

Ala Met Val Leu Trp Val Phe Phe Leu Val Thr Leu Thr Leu Ser Ser 1 5 10 15

Gly Ser His Gly Ser Leu Pro Ser Gln Pro Leu Arg Ile Pro Arg Tyr
20 25 30

Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln Leu 35 40 45

Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Asn Arg Gln Gln Gly Glu 50 55 60

Arg Asn Gln Glu Gln Gly Ala Ala Cys 65 70

<210> 30

<211> 222

<212> DNA

<213> artificial sequence

<220>

<223> Nucleic acid sequence for an original bovine GHRH sequence

ccatggtgct ctgggtgttc ttcctcgtga ccctcaccct cagcagcggc tcccacggtt

<400> 30

ccctgccttc ccagcctctc aggattccac ggtacgccga cgccatcttc accaacagct 120
accggaaggt gctgggccag ctgtccgccc ggaagctgct gcaggacatc atgaacaggc 180
agcagggcga gagaaaccag gagcagggcg cctgataagc tt 222

60

<210> 31

<211> 71

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for an original bovine GHRH sequence

<400> 31

Met Val Leu Trp Val Phe Phe Leu Val Thr Leu Thr Leu Ser Ser Gly

1 5 10 15

Ser His Gly Ser Leu Pro Ser Gln Pro Leu Arg Ile Pro Arg Tyr Ala 20 25 30

Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln Leu Ser 35 40 45

Ala Arg Lys Leu Leu Gln Asp Ile Met Asn Arg Gln Gln Gly Glu Arg 50 55 60

Asn Gln Glu Gln Gly Ala Ala 65 70

<210> 32

<211> 73

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for an analog ovine GHRH sequence

<400> 32

Ala Met Val Leu Trp Val Phe Phe Leu Val Thr Leu Thr Leu Ser Ser 1 5 10 15

Gly Ser His Gly Ser Leu Pro Ser Gln Pro Leu Arg Ile Pro Arg Tyr
20 25 30

Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Ile Leu Gly Gln Leu 40 35 Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Asn Arg Gln Gln Gly Glu 55 60 50 Arg Asn Gln Glu Gln Gly Ala Ala Cys 65 70 <210> 33 <211> 222 <212> DNA <213> artificial sequence <220> <223> Nucleic acid sequence for an original ovine GHRH sequence <400> 33 ccatggtgct ctgggtgttc ttcctcgtga ccctcaccct cagcagcggc tcccacggtt 60 ccctgccttc ccagcctctc aggattccac ggtacgccga cgccatcttc accaacagct 120 accggaagat cctgggccag ctgtccgccc ggaagctgct gcaggacatc atgaacaggc 180 222 agcagggcga gagaaaccag gagcagggcg cctgataagc tt <210> 34 <211> 71 <212> PRT <213> artificial sequence

<223> Amino acid sequence for an original ovine GHRH sequence

<220>

<400> 34

Met Val Leu Trp Val Phe Phe Leu Val Thr Leu Thr Leu Ser Ser Gly

1 5 10 15

Ser His Gly Ser Leu Pro Ser Gln Pro Leu Arg Ile Pro Arg Tyr Ala 20 25 30

Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Ile Leu Gly Gln Leu Ser 35 40 45

Ala Arg Lys Leu Leu Gln Asp Ile Met Asn Arg Gln Gln Gly Glu Arg 50 55 60

Asn Gln Glu Gln Gly Ala Ala 65 70

<210> 35

<211> 234

<212> DNA

<213> artificial sequence

<220>

<223> Nucleic acid sequence for an analog chicken GHRH sequence

<400> 35

gccatggccc tgtgggtgtt ctttgtgctg ctgaccctga cctccggaag ccactgcagc 60
ctgccaccca gcccaccctt ccgcgtcagg cgccacgccg acggcatctt cagcaaggcc 120
taccgcaagc tcctgggcca gctgagcgca cgcaactacc tgcacagcct gatggccaag 180
cgcgtgggca gcggactggg agacgaggcc gagcccctga gctgataagc ttgc 234

<210> 36

<211> 76

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for an analog chicken GHRH sequence

<400> 36

Ala Met Ala Leu Trp Val Phe Phe Val Leu Leu Thr Leu Thr Ser Gly

1 5 10 15

Ser His Cys Ser Leu Pro Pro Ser Pro Pro Phe Arg Val Arg Arg His
20 25 30

Ala Asp Gly Ile Phe Ser Lys Ala Tyr Arg Lys Leu Leu Gly Gln Leu 35 40 45

Ser Ala Arg Asn Tyr Leu His Ser Leu Met Ala Lys Arg Val Gly Ser 50 55 60

Gly Leu Gly Asp Glu Ala Glu Pro Leu Ser Ala Cys
65 70 75

<210> 37

<211> 231

<212> DNA

<213> artificial sequence

<220>

<223> Nucleic acid sequence for an original chicken GHRH sequence
<400> 37
ccatggcact ctgggtgttc tttgtgctcc tcaccctcac cagtggctcc cactgctcac
60
tgcccccctc acctcccttc agggtgcgc ggcacgccga tgggatcttc agcaaagcct
120
acaggaaact cctgggccag ctgtccgcaa gaaattacct gcactccctg atggccaagc
180

231

<210> 38

<211> 74

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for an original chicken GHRH sequence

gggtcggcag cggcctgggg gacgaggcgg aaccgctcag ctgataagct t

<400> 38

Met Ala Leu Trp Val Phe Phe Val Leu Leu Thr Leu Thr Ser Gly Ser 1 5 10 15

His Cys Ser Leu Pro Pro Ser Pro Pro Phe Arg Val Arg Arg His Ala 20 25 30

Asp Gly Ile Phe Ser Lys Ala Tyr Arg Lys Leu Leu Gly Gln Leu Ser 35 40 45

Ala Arg Asn Tyr Leu His Ser Leu Met Ala Lys Arg Val Gly Ser Gly 50 55 . 60

Leu Gly Asp Glu Ala Glu Pro Leu Ser Ala 70 65 <210> 39 <211> 40 <212> PRT <213> artificial sequence <220> <223> Amino acid sequence for GHRH sequence wt-GHRH <400> 39 Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln 15 10 Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly 25 30 20 Glu Arg Asn Gln Glu Gln Gly Ala 35 40 <210> 40 <211> 40 <212> PRT <213> artificial sequence

<220>

<223> Amino acid sequence for GHRH sequence HV-GHRH

<400> 40

His Val Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Ala Gln

1 5 10 15

-41-

4431793v.1

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Leu Asn Arg Gln Gln Gly
20 25 30

Glu Arg Asn Gln Glu Gln Gly Ala 35 40

<210> 41

<211> 40

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for GHRH sequence TI-GHRH

<400> 41

Tyr Ile Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Ala Gln

1 5 10 15

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Leu Asn Arg Gln Gly 20 25 30 $^{\circ}$

Glu Arg Asn Gln Glu Gln Gly Ala 35 40

<210> 42

<211> 40

<212> PRT

<213> artificial sequence

<220>

-42-

<223> Amino acid sequence for GHRH sequence TV-GHRH

<400> 42

Tyr Val Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Ala Gln 10 15

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Leu Asn Arg Gln Gln Gly 25 30 20

Glu Arg Asn Gln Glu Gln Gly Ala 35 40

<210> 43

<211> 40

<212> PRT

<213> artificial sequence

<220>

<223> Amino acid sequence for GHRH sequence 15/27/28-GHRH

<400> 43

Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Ala Gln 15 10

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Leu Asn Arg Gln Gln Gly 20 25 30

Glu Arg Asn Gln Glu Gln Gly Ala 35

40